

AP20 Rec'd PCT/PTO 22 JUN 2006

SEQUENCE LISTING

<110> CropDesign N.V.

<120> Plants having increased yield and method for making the same

<130> CD-106-PCT

<150> US 60/532,287

<151> 2003-12-22

<160> 5

<170> PatentIn version 3.3

<210> 1

<211> 1311

<212> DNA

<213> Arabidopsis thaliana

<220>

<221> misc_feature

<223> A variant of the coding sequence of the sequence deposited under accession number NM_121168 contains a G instead of C on position 851 and a T instead of C on position 1295

<400> 1

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gtatcaatac	ctccaacaaa	accttctttt	aaacagcaaa	agagacgtgc	agtacttaag	180
gatgtgagta	ataccctctgc	agatattattt	tattcagaac	ttcgaaaggg	aggcaacatc	240
aaggcaaca	gaaaatgtct	aaaagagcct	aaaaaagcag	caaaggaagg	tgctaacagt	300
gccatggata	ttctggtaga	tatgcataca	gaaaaatcaa	aatttagcaga	agatttgccc	360
aatgatcaga	tggtctgaa	ccaagatgtc	tctttttcaa	actttaaaaga	tgaagaaaatt	420
actcgacaa	aagaatgtgg	atcagggtgc	atggagttac	ttcaagttgt	agatattgtat	480
tccaaacgtcg	aagatccaca	gtgttgcagc	ttgtatgctg	ctgatataata	tgacaacata	540
catgttgcag	agcttcaaca	acgacccttg	gctaattttt	tggagcttgc	gcagcgagat	600
atcgaccac	acatgagaaaa	gattctgatt	gactggcttg	tagaagtttc	tgacgactac	660
aagctgggtc	cagatacgct	ttaccttaca	gtgaatctta	tcgaccgggtt	tctgtccaa	720
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aaatatgaag	agctttccgc	accaggggtt	gaggagttt	gcttcattac	ggccaacaca	840
tacacaagac	cagaagtgc	gagcatggag	attccaaattt	taaattttgt	gcacttttgc	900
ttatcggttc	ctaccaccaa	aacatttctg	aggcggttca	ttaaagcagc	tcaagcttcg	960
tacaagggtc	ctttcattga	actggagttat	ttagcaact	atctcgccga	attgacactg	1020
gtggaatata	gtttcctaag	gttcctgcca	tcactaattt	ctgcttcagc	tgttttccca	1080
gccccgatgga	cactcgacca	aactgaccat	ccttggaaacc	ctactctgca	acactacacc	1140
agatatgagg	tagctgagct	gaagaacaca	gttctcgcca	tggaggactt	gcagctcaac	1200
accagtggct	gtactctcg	tgccacccgt	gagaaataca	accaacccaa	gtttaagagc	1260
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<210> 2

<211> 436

<212> PRT

<213> Arabidopsis thaliana

<220>

<221> MISC_FEATURE

<223> A variant of the sequence deposited under accession number NP_568248 contains an arginine instead of a proline on position

284 and a phenylalanine instead of a serine on position 432

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				20				25							30
Ala	Lys	Lys	Ala	Met	Gly	Arg	Gly	Val	Ser	Ile	Pro	Pro	Thr	Lys	Pro
				35				40							45
Ser	Phe	Lys	Gln	Gln	Lys	Arg	Arg	Ala	Val	Leu	Lys	Asp	Val	Ser	Asn
				50				55				60			
Thr	Ser	Ala	Asp	Ile	Ile	Tyr	Ser	Glu	Leu	Arg	Lys	Gly	Gly	Asn	Ile
				65				70							80
Lys	Ala	Asn	Arg	Lys	Cys	Leu	Lys	Glu	Pro	Lys	Lys	Ala	Ala	Lys	Glu
				85				90							95
Gly	Ala	Asn	Ser	Ala	Met	Asp	Ile	Leu	Val	Asp	Met	His	Thr	Glu	Lys
				100				105							110
Ser	Lys	Leu	Ala	Glu	Asp	Leu	Ser	Lys	Ile	Arg	Met	Ala	Glu	Ala	Gln
				115				120							125
Asp	Val	Ser	Leu	Ser	Asn	Phe	Lys	Asp	Glu	Glu	Ile	Thr	Glu	Gln	Gln
				130				135							140
Glu	Asp	Gly	Ser	Gly	Val	Met	Glu	Leu	Leu	Gln	Val	Val	Asp	Ile	Asp
				145				150							160
Ser	Asn	Val	Glu	Asp	Pro	Gln	Cys	Cys	Ser	Leu	Tyr	Ala	Ala	Asp	Ile
				165				170							175
Tyr	Asp	Asn	Ile	His	Val	Ala	Glu	Leu	Gln	Gln	Arg	Pro	Leu	Ala	Asn
				180				185							190
Tyr	Met	Glu	Leu	Val	Gln	Arg	Asp	Ile	Asp	Pro	Asp	Met	Arg	Lys	Ile
				195				200							205
Leu	Ile	Asp	Trp	Leu	Val	Glu	Val	Ser	Asp	Asp	Tyr	Lys	Leu	Val	Pro
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Asp	Thr	Leu	Tyr	Leu	Thr	Val	Asn	Leu	Ile	Asp	Arg	Phe	Leu	Ser	Asn
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Ser	Tyr	Ile	Glu	Arg	Gln	Arg	Leu	Gln	Leu	Leu	Gly	Val	Ser	Cys	Met
				245				250							255
Leu	Ile	Ala	Ser	Lys	Tyr	Glu	Glu	Leu	Ser	Ala	Pro	Gly	Val	Glu	Glu
				260				265							270
Phe	Cys	Phe	Ile	Thr	Ala	Asn	Thr	Tyr	Thr	Arg	Pro	Glu	Val	Leu	Ser
				275				280							285
Met	Glu	Ile	Gln	Ile	Leu	Asn	Phe	Val	His	Phe	Arg	Leu	Ser	Val	Pro
				290				295							300

Thr Thr Lys Thr Phe Leu Arg Arg Phe Ile Lys Ala Ala Gln Ala Ser
 305 310 315 320

Tyr Lys Val Pro Phe Ile Glu Leu Glu Tyr Leu Ala Asn Tyr Leu Ala
 325 330 335

Glu Leu Thr Leu Val Glu Tyr Ser Phe Leu Arg Phe Leu Pro Ser Leu
 340 345 350

Ile Ala Ala Ser Ala Val Phe Leu Ala Arg Trp Thr Leu Asp Gln Thr
 355 360 365

Asp His Pro Trp Asn Pro Thr Leu Gln His Tyr Thr Arg Tyr Glu Val
 370 375 380

Ala Glu Leu Lys Asn Thr Val Leu Ala Met Glu Asp Leu Gln Leu Asn
 385 390 395 400

Thr Ser Gly Cys Thr Leu Ala Ala Thr Arg Glu Lys Tyr Asn Gln Pro
 405 410 415

Lys Phe Lys Ser Val Ala Lys Leu Thr Ser Pro Lys Arg Val Thr Ser
 420 425 430

Leu Phe Ser Arg
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<210> 3

<211> 654

<212> DNA

<213> Oryza sativa

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aaacaagagt gtcaatggaa caatggaaaac catatgacat actataattt tgTTTTatt	240
attgaaatttataattcaa agagaataaa tccacatagc cgtaaaagttc tacatgtgg	300
gcattaccaa aatatataaa gcttacaaaaa catgacaaggc tttagttgaa aaatttgcatt	360
ccttattcaca ttgacacata aagtggatgtga tgagtctaa tattatttc tttgtcaccc	420
atcatgtata tatgatagcc acaaaagttag tttgtatgtat atatcaaaga acatttttag	480
gtgcacctaa cagaatatcc aaataatatg actcaacttag atcataatag agcatcaagt	540
aaaactaaca ctctaaagca accgatggaa aagcatctat aaatagacaa gcacaatgaa	600
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<210> 4

<211> 56

<212> DNA

<213> Artificial sequence

<220>

<223> primer PRM582

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<211> 52

<212> DNA
<213> Artificial sequence

<220>
<223> primer PRM583

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52